

UniprotKB ID	Entry name	organism	full name	oglcna score	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma meml
Q9WTP0	E41L1_RAT	Rattus norvegicus	Band 4.1-like protein 1	41.439696	NaN	T30;S75;T79;T343;S378;S430;S437;S461;S466;T475;S510;S540;S541;S544;S546;T550;S564;S578;T580;S639;S648;S650;S665;S666;S671;S677;S684;T685;S721;S782;S868;S583;S587;S669;S742;S766;S968;S1092;S1223	18683930;15340146;17496889;19322778	MTTETGPDSEVKKAEETPQQPEAA AAVTTPTVTPAGHSHPETNSNEKHLT QQDTRPAEQSLDMEEKDYCEADGL SERTTPSKAQKSPQKIAKKFKSATCR VTLLDASEYECEVEKHKHGRGQVFLDL VCEHLNLLLEKDYFGLTFCDADSQKN WLDPSKEIKKQIRSSPWNFAFTVKFY PPDPAQLTEDITRYLCLQLRADIITG RLPCSFVTHALLGSYAVQAEGLGDHD TEEHVGNVSELRFAPNQTRELEER IMELHKTYRGMTPGEAEIHFLENAK KLSMYGVDLHHAKDSEGDIMLGVC ANGLLIYRDLRINRFAWPKILKISYK RSNFYKIRPGEYEQFESTIGFKLPN HRSKRLLWKVCIHHTFFRLVSPPEP PPKGFVLMGSKFRYSGRTOAQTRQA SALIDRPAPFFERSSSKRYTMSRSLD GAEFSRPASVSENHDAGPDGDKRE DDAESGRRSEAEEGEVTRPTKIKEL KPEQETTPRHKQEFLDKPEDVLLKH QASINELKRTLKEPNSKLHRRDRDW ERERRLPSSPASPSPKCTPEKASERA GLREGSEEKVKPPRPRAPESDTGDE DQDQERDAVFLKDNHLAIERKCSSI TVSSTSSLEAEVDFTVIGDYHGGAFE DFSRLPELDRDKSDSETGLVFAR DLKGPSSQEDES GGIEDSPDRGACS TPELPQFESVKAETMTVSSLAIRKKI EPEAMLQSRVSTADSTQVDGGAPAA KDFMTTPPCITTEITSTTMENSLKSG KGAAAMIPGPQTVATEIRLSPIIGKD VLTSTYGATAETLSTSTTHVTKTVK GGFSETRIEKRIITGDEDVDQDQAL ALAIKEAKLQHPDMLVTKAVVYRET DPSPEERDKKPQES	None	None	None	None	None	None	None	

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